

Before going **BIG** - seize opportunities for **small** data: multistate models to improve decision-making and effect quantification in clinical trials

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Who



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Multistate models in clinical trials

Where



RESEARCH PAPER | Full Access |

A multistate model for early decision-making in oncology

Ulrich Beyer , David Dejardin, Matthias Meller, Kaspar Rufibach, Hans Ulrich Burger

First published: 16 July 2019 | <https://doi.org/10.1002/bimj.201800250>

Simulated dataset + comprehensive R code.

**How do we typically decide whether
to move an oncology molecule
into Phase 3?**

Decision-making in early oncology development

- 1 Small single-arm trial for **experimental** drug (e.g. $n = 40$).
- 2 Response proportion, duration of response.
- 3 Compare to “corresponding” quantities from literature for **control** treatment.

But:

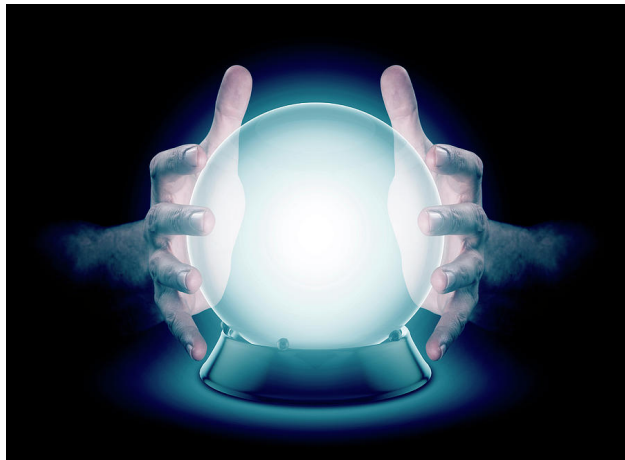
- **P(wrong decision)** may be high.
- Primary endpoint in Phase 3: **Overall survival**.

Proposal:

Decide in early phase based on OS prediction.

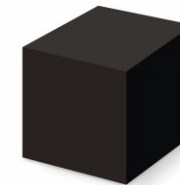
Decrease P(wrong decision).

Prediction?



Prediction?

Blackbox ML algorithm using big data?



Transparent multistate model with historical borrowing.

Challenges and proposal

Challenges:

- 1 Response-type endpoint?
- 2 Surrogacy? **Poor** in many indications.
- 3 Immunotherapy (CIT): no effect on response, relevant OS effect.
- 4 **Non-randomized** comparison \Rightarrow confounding.

Proposal: Base decision-making on **OS prediction from multistate model**.

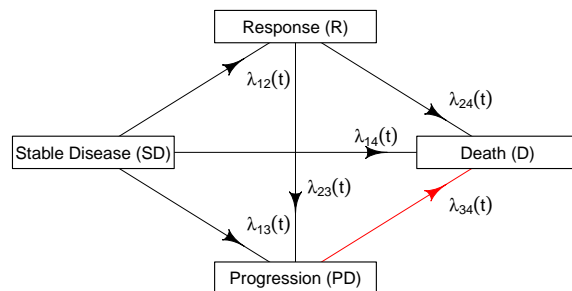
- 1 **Predicted survival function for experimental arm.**
- 2 Combine S_{exp} with S_{control} to get **predicted OS HR**.
- 3 Experimental drug might act on certain transitions only \Rightarrow not captured through simple modelling of OS. Potential **efficiency gain!**
- 4 **Propensity scoring.**

Idealized scenario: Retrospective data from Phase 3 RCTs.

Long-term follow-up in both arms.

Randomization \Rightarrow no confounding.

Multistate model for early decision-making



- Follow-up of patient until **PD or death without PD**.
- Post-progression hazard λ_{34} : **borrowing** from historical data.
- Transitions $SD \rightarrow D, R \rightarrow D$ rare, hazards \approx same in both arms.

Predicted survival function in experimental arm, S_{exp}

Compute transition probabilities for each transition.

$$S_{\text{exp}}(t) = 1 - \left(P_{SD \rightarrow D}(0, t) + P_{SD \rightarrow \text{PD} \rightarrow D}(0, t) + P_{SD \rightarrow R \rightarrow D}(0, t) + P_{SD \rightarrow R \rightarrow \text{PD} \rightarrow D}(0, t) \right).$$

λ_{34} corresponding to **PD \rightarrow D** transition borrowed from historical data.

Historical borrowing for λ_{34}

Experimental treatment expected to provide benefit **beyond PD**?

No:

- E.g. chemotherapy or antibody-dependent cellular cytotoxicity.
- **Plug-in** hazard function estimate from historical control.
- No post-PD information required for experimental arm.

Yes:

- E.g. chemoimmunotherapy.
- Estimate post-PD hazard ratio assuming **proportionality**.
- How much post-PD deaths needed in experimental arm to reliably **estimate post-PD HR**?

Benefit beyond PD: Oak

Oak

Previously treated non-small-cell lung cancer.

Rittmeyer et al. (2017).

	Atezolizumab	Chemotherapy	Hazard ratio
Effect post-PD	expected	not expected	
Objective Response	58 (13.6%)	57 (13.4%)	
Duration of Response	26.3 (10 - ∞)	6.2 (4.9 - 7.6)	
Overall Survival			0.73 (0.62, 0.87)

**If this were early phase data -
would you initiate Phase 3?**

**Competitors used this
mechanism of action.**

OS prediction when post-PD hazards assumed proportional

Random variable:

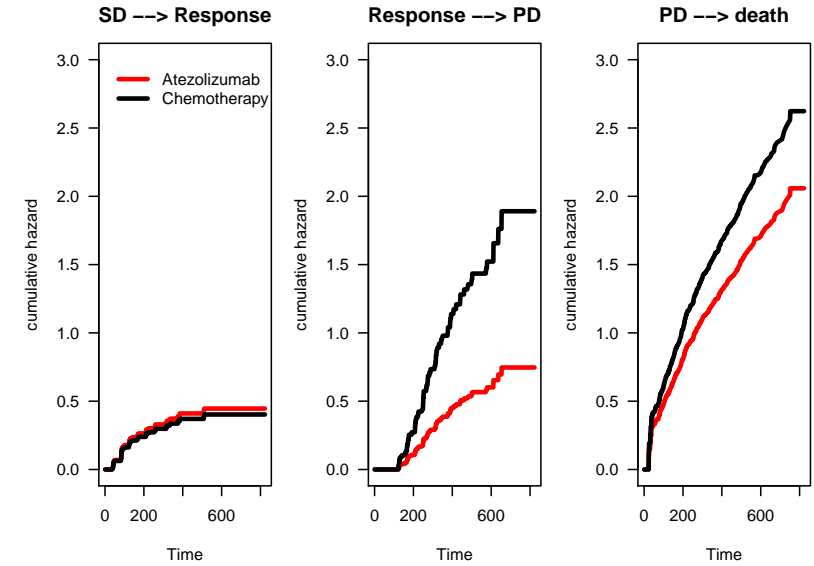
$$Z = \begin{cases} 0 & \text{if patient in control,} \\ 1 & \text{if in experimental group.} \end{cases}$$

$$\lambda_{34}(t|Z) = \lambda_{34,0}(t) \exp(\beta_{34}Z)$$

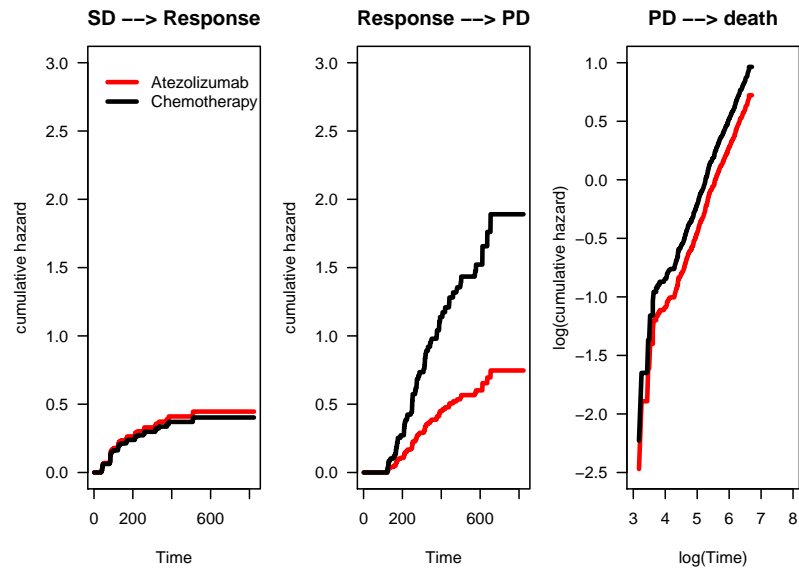
Baseline hazard $\lambda_{34,0}$ **estimated from both arms combined.**

Post-progression hazard ratio β_{34} ?

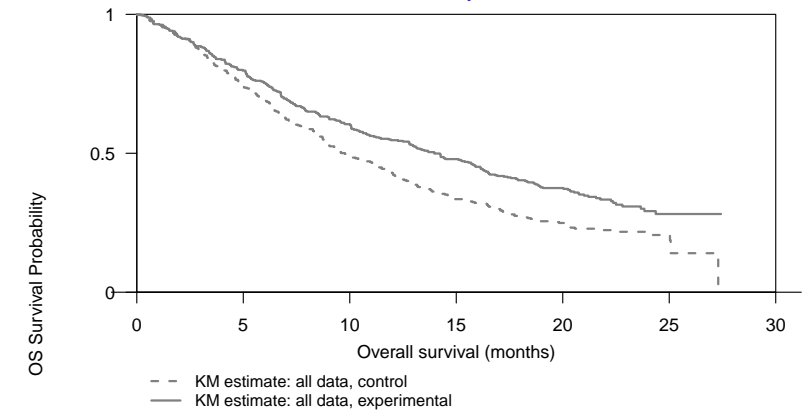
Oak: raw cumulative hazard estimates (of interest)



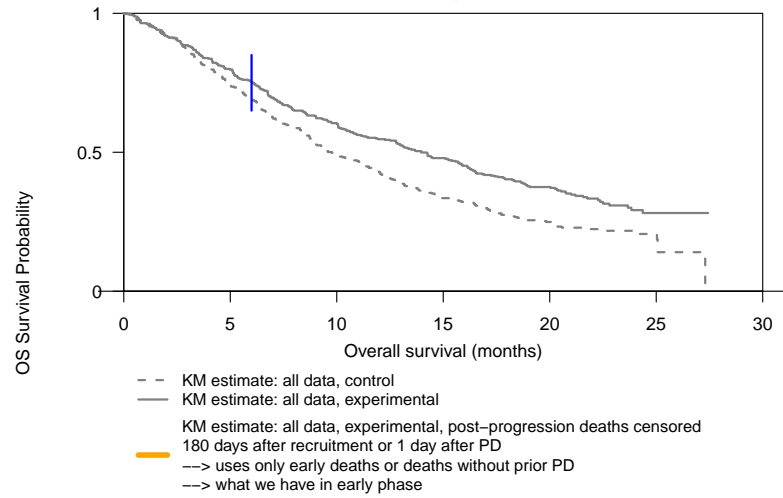
Oak: raw cumulative hazard estimates (of interest)



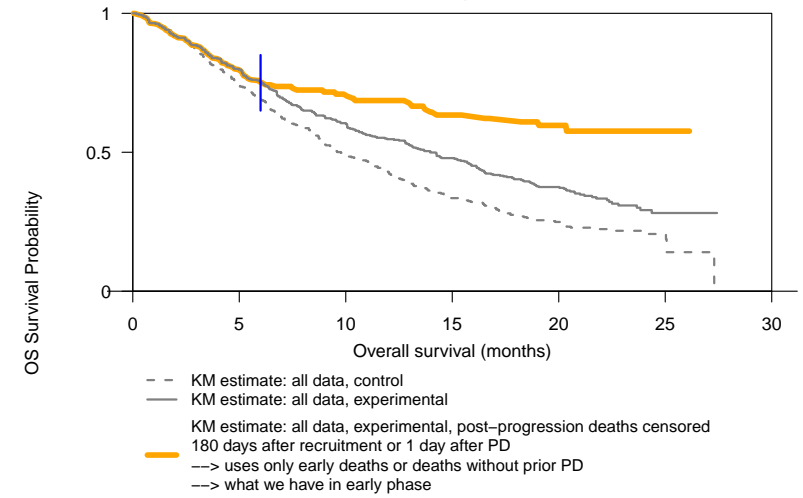
Oak: estimates / predictions of S_{exp}



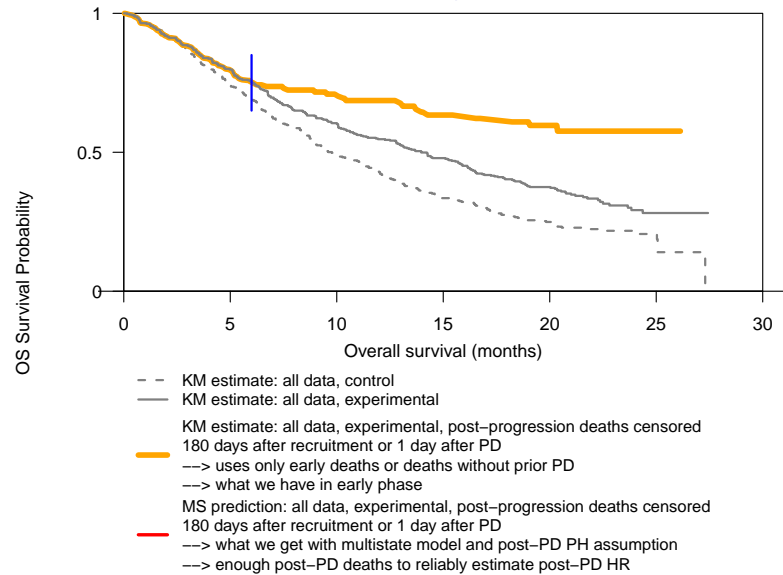
Oak: estimates / predictions of S_{exp}



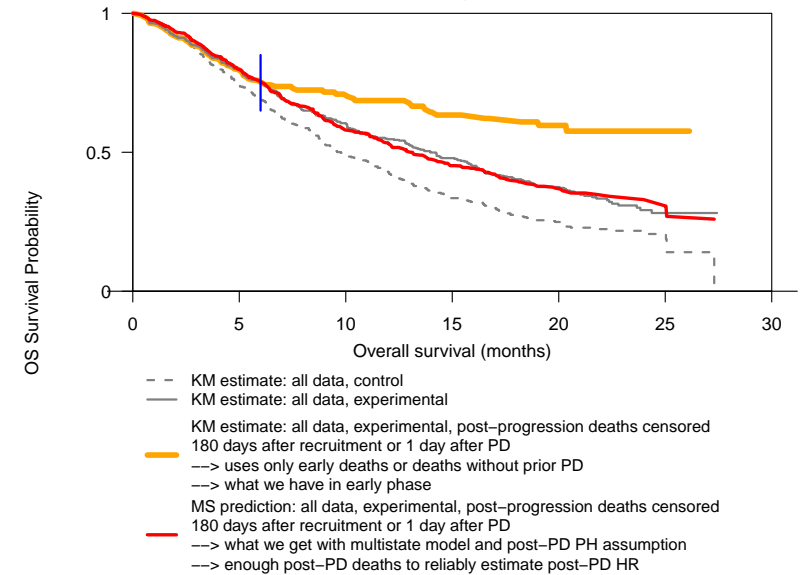
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Oak: estimates / predictions of S_{exp}



Early phase decision based on multistate prediction:

P(wrong decision)?

OS prediction from mimicked early phase data

Historical control: Oak control arm data.

False-positive decision: Sample early phase trial from Oak control arm.

False-negative decision: Sample early phase trial from Oak experimental arm.

Sample early phase trial:

- 40 patients,
- 6 months uniform recruitment,
- analysis 15 months after first patient entered,
- censor post-PD follow-up **one day after PD**,
- estimate $\lambda_{12}, \lambda_{13}, \lambda_{14}, \lambda_{23}, \lambda_{24}$ from this data.

Cox regression for post-PD transition $\Rightarrow \hat{\lambda}_{34}(t|Z)$.

Compute prediction of S_{exp} .

OS HR prediction based on early phase trial

Approximate HR by fitting exponential distribution to both arms $\Rightarrow \widehat{HR}$.

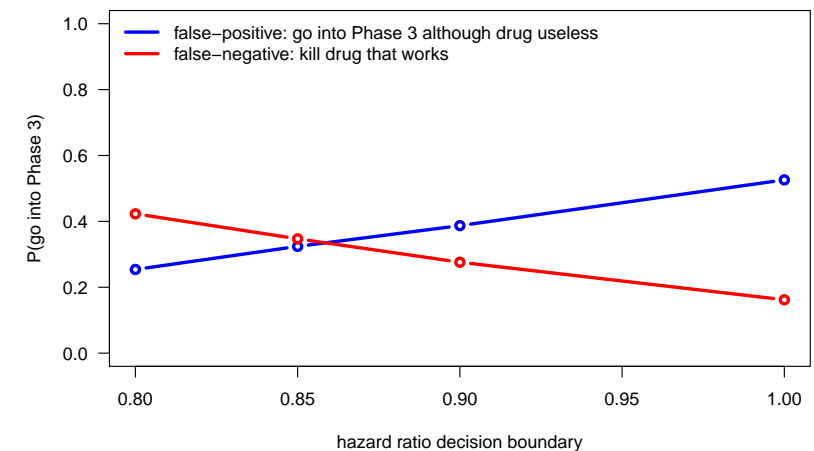
Decision to move to Phase 3: $\widehat{HR} \leq \text{boundary} \in \{0.80, 0.85, 0.90, 1.00\}$.

Repeat 1000 times.

Resampling \Rightarrow **quantification of uncertainty**.

Oak: P(wrong decision)

P(go into Phase 3) = P(approximated HR \leq boundary)



How many post-PD deaths to estimate HR of PD \rightarrow death transition?

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Ask during Q&A.

Conclusions for early-decision making proposal

Conclusions

Early phase decision-making based on **multistate OS prediction**:

- Assumption on λ_{34} \Rightarrow need to understand **disease and treatment**.
- **Avoids difficulty in interpretation of response-type endpoints**.
- Feasibility assessed in **idealized scenario**.
- Recommendation **how much post-PD follow-up** needed to estimate β_{34} .
- Needs **long-term individual-patient** data in control arm!

What about confounding?

Real-world data as historical control.

Combine proposal with propensity scoring.

Immunotherapy:
1) no difference in PFS,
2) non-proportional hazards for OS.

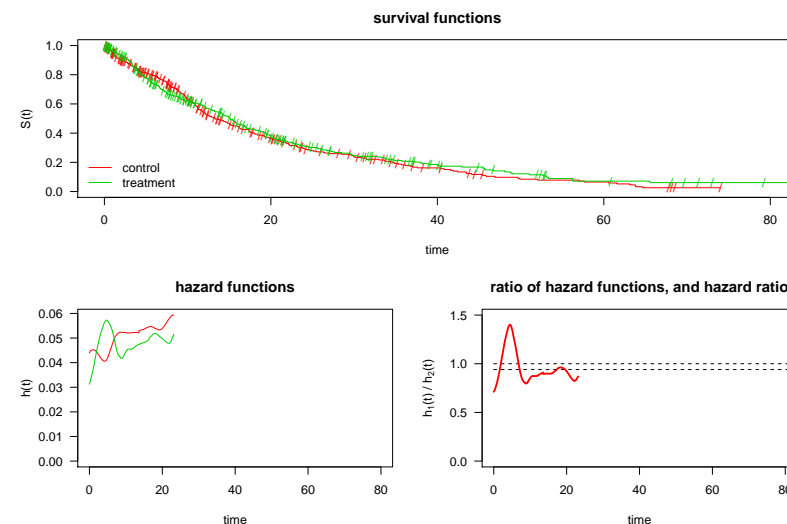
How to quantify effect?

A fictional clinical trial

Simulated clinical trial:

- 1:1 randomized, 400 and 400 patients per arm.
- No administrative censoring, but drop-out.

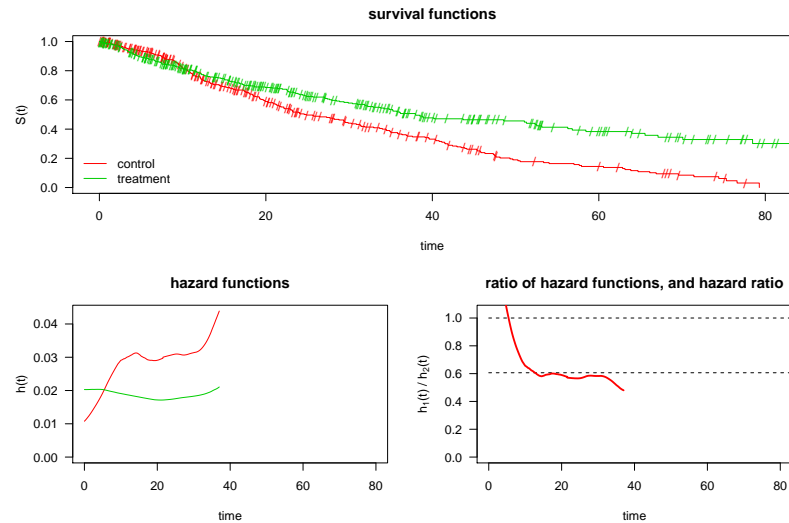
PFS for simulated clinical trial



- Estimated hazard ratio: 0.94, 95% confidence interval [0.80, 1.11].

- Hypothesis test for PH: $p = 0.24$.

OS for simulated clinical trial



- Estimated hazard ratio: 0.61, 95% confidence interval [0.50, 0.74].
- Hypothesis test for PH: $p < 0.0001$.

Summarize treatment effect

Non-proportional hazards for OS. How to summarize effect of treatment?

Data was generated according to:

Transition	Control arm	Treatment arm
$0 \rightarrow 1$	$\lambda_{01}^c = \log(2)/25$	$\lambda_{01}^t = \lambda_{01}^c \cdot \mathbf{1}$
$0 \rightarrow 2$	$\lambda_{02}^c = \log(2)/30$	$\lambda_{02}^t = \lambda_{02}^c \cdot \mathbf{0.8}$
$1 \rightarrow 2$	$\lambda_{12}^c = \log(2)/15$	$\lambda_{12}^t = \lambda_{12}^c \cdot \mathbf{0.4}$

	coef	HR = exp(coef)	95% CI	p-value
transition event-free \rightarrow PD	-0.04	0.96	[0.77, 1.19]	0.72
transition event-free \rightarrow death	-0.09	0.91	[0.70, 1.18]	0.49
transition PD \rightarrow death	-1.09	0.34	[0.24, 0.46]	< 0.0001

Gaschler-Markefski et al. (2014).

Conclusions

Multistate models

Multistate models useful:

- Canonical **extension of survival analysis**.
- Get more **insight** in how disease and drug work.
- **Prediction** in well-specified, as opposed to black-box, model.
- **Jointly** model three key oncology endpoints: response, PFS, OS.
- Applications by no means restricted to oncology!

Many potential applications:

- Improved **early stage decision-making** \Rightarrow Beyer et al. (2019).
- Improved **communication** of effect and optimized **sample size** computation.
- Bivariate modelling of PFS and OS to help inform **surrogacy** questions \Rightarrow Meller et al. (2019).

Big vs. small data

Often, information removed/altered in **small data**:

- (Artificial) response categories instead of actual measurements: **dichotomization**,
- response proportions only: **ignoring the dynamics between states**,
- complicated subsets, e.g. those that respond only: **selection bias**,
- effect quantification in **one** number where biological process might suggest few numbers,
- ...

Thank you for your attention

Maximize information from **small** data. AND look at **BIG** data.

Biostatisticians ideally placed to contribute to this!

References I

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Doing now what patients need next

R version and packages used to generate these slides:

R version: R version 3.6.0 (2019-04-26)

Base packages: stats / graphics / grDevices / utils / datasets / methods / base

Other packages: nls2 / proto / diagram / shape / ggplot2 / rocheBCE / muhaz / flexsurv / reporttools / xtable / mstate / etm / dplyr / mvna / prodlim / biostatKR / survival

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